



SEQ ID NO: 1

SPAPPACDLRVLSKLLRDSHVLHSRLSQCEVHPLPTPVLLPAVDFSLGEWKTQMEETKAQDI  
LGAVTLLLEGVMAARGQLGPTCLSSLLGQLSGQVRLLLGALQSLGTLPPQGRTTAHKDPNA  
IFLSFQHLLRGKVRFLMLVGGSTLCVRRAPPTTAVPS

**FIG.\_1**

SEQ ID NO:2

QACKKHELYVSFRDLGWQDWIIAPEGYAAYYCEGECAPFLNSYMNATNHAIVQTLVHFINPET  
VPKPCCAPTQLNAISVLYFDDSSNVILKKYRNMVVRACGCH

**FIG.\_2**

SEQ ID NO:3

MSYNLLGFLQRSSNFQCQKLLWQLNGRLEYCLKDRMNFDIPEEIKQLQQFQKEDAALTIYEML  
QNI FAIFRQDSSSTGWN ETIVENLLANVYHQINHLKTVLEEKLEKEDFTRGKLMSSLHLKRY  
GRILHYLKAKEYSHCAWTIVRVEILRN FYFINRLTG YLRN

**FIG.\_3**

SEQ ID NO:4

MAFTEHSPLTPHRRDLCSRSIWLARKIRSDLTALTESYVKHQGLNKNINLDSADGMPVASTDQ  
WSELTEAERLQENLQAYRTFHVLLARLLEDQQVHFTPTGDFHQAIHTLLLQVAAFAYQIEEL  
MILLEYKIPRNEADGMPINVG DGGLFEKKLWGLKVLQELSQWTVRSIHDLRFISSHQGTGIPAR  
GSHYIANNKKM

**FIG.\_4**

